

## STIC Biotechnology Systems Branch

### RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/031,639A  
Source: I#W16  
Date Processed by STIC: 3-19-05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - cPAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):  
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



IFW16

## RAW SEQUENCE LISTING

DATE: 03/19/2005

PATENT APPLICATION: US/10/031,639A

TIME: 10:33:53

Input Set : A:\SEQLIST.txt

Output Set: N:\CRF4\03192005\J031639A.raw

4 <110> APPLICANT: CELL GENESYS, INC.  
 6 <120> TITLE OF INVENTION: METHODS AND MEANS FOR PRODUCING HIGH  
 7 TITER, SAFE, RECOMBINANT LENTIVIRUS VECTORS  
 10 <130> FILE REFERENCE: G&C 131.3USWO  
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/031,639A  
 C--> 12 <141> CURRENT FILING DATE: 2002-05-09  
 12 <150> PRIOR APPLICATION NUMBER: 10/031,639  
 13 <151> PRIOR FILING DATE: 2001-10-29  
 15 <160> NUMBER OF SEQ ID NOS: 52  
 17 <170> SOFTWARE: FastSEQ for Windows Version 4.0

## ERRORED SEQUENCES

584 <210> SEQ ID NO: 52  
 585 <211> LENGTH: 21  
 586 <212> TYPE: DNA  
 587 <213> ORGANISM: Artificial Sequence  
 589 <220> FEATURE:  
 590 <223> OTHER INFORMATION: linker  
 592 <400> SEQUENCE: 52  
 593 gatcaagcct actcgacccg g  
 E--> 605 1

Does Not Comply  
Corrected Diskette Needed

(pg. 12)

21

P/S delete

<210> 31  
<211> 10  
<212> DNA  
<213> Artificial sequence  
~~<220>~~  
~~<223>~~  
<400> 31  
ggccattgac

Insert

pls explain  
Source of genetic  
material. See  
error explanation  
on page 3.

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/031,639A

DATE: 03/19/2005  
TIME: 10:33:54

Input Set : A:\SEQLIST.txt

Output Set: N:\CRF4\03192005\J031639A.raw

Use of <220> Feature (NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.

Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:31

*ERROR EXPLANATION: ✓*

## VERIFICATION SUMMARY

DATE: 03/19/2005

PATENT APPLICATION: US/10/031,639A

TIME: 10:33:54

Input Set : A:\SEQLIST.txt

Output Set: N:\CRF4\03192005\J031639A.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No  
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:358 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:31  
L:360 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:31, <213>  
ORGANISM:Artificial Sequence  
L:360 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:31, <213>  
ORGANISM:Artificial Sequence  
L:360 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:31,Line#:360  
L:605 M:254 E: No. of Bases conflict, this line has no nucleotides.